

SEQUENCE LISTING



<110> DENEFLÉ, Patrice
 ROSIER-MONTUS, Marie-Francoise
 PRADES, Catherine
 ARNOULD-REGUIGNE, Isabelle
 DUVERGER, Nicolas
 ALLIKMETS, Rando
 DEAN, Michael

<120> NUCLEIC ACIDS OF THE HUMAN ABCA5, ABCA6, ABCA9, AND ABCA10 GENES, VECTORS
 CONTAINING SUCH NUCLEIC ACIDS, AND USES THEREOF

<130> ABCA5, 6, 9, 10

<140> US 10/005,338

<141> 2001-12-07

<150> US 60/263,231

<151> 2001-01-23

<150> FR 00403440.1

<151> 2000-12-07

<160> 217

<170> PatentIn Ver. 2.1

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| atttcaggcc | tctggccttc | agcatactgt | tgtggacagg | ctctgggtga | cattccatta | 3840 |
| tacttcttga | ttctcttttc | aatacattta | atttactact | tcataattct | gggattccag | 3900 |
| ctttcatggg | aactcatgtt | tgttttgggt | gtatgcataa | ttggttgtgc | agtttctctt | 3960 |
| atattcctca | catatgtgct | ttcattcatc | tttcgcaagt | ggagaaaaaa | taatggcttt | 4020 |
| tgggtctttg | gcttttttat | tatcttaata | tgtgtatcca | caattatggg | atcaactcaa | 4080 |
| tatgaaaaac | tcaacttaat | tttgtgcatg | attttcatac | cttccttcac | tttgcctggg | 4140 |
| tatgtcatgt | tattgatcca | gctcgacttt | atgagaaact | tggacagtct | ggacaataga | 4200 |
| ataaatgaag | tcaataaaaac | cattctttta | acaaccttaa | taccatacct | tcagagtgtt | 4260 |
| attttctttt | ttgtcataag | gtgtctggaa | atgaagtatg | gaaatgaaat | aatgaataaa | 4320 |
| gacccagttt | tcagaatctc | tccacggagt | agagaaactc | atcccaatcc | ggaagagccc | 4380 |
| gaagaagaag | atgaagatgt | tcaagctgaa | agagtccaag | cagcaaatgc | actcactgct | 4440 |
| ccaaacttgg | aggaggaacc | agtcataact | gcaagctgtt | tacacaagga | atattatgag | 4500 |
| acaaagaaaa | gttgcttttc | aacaagaaag | aagaaaaatg | ccatcagaaa | tgtttctttt | 4560 |
| tgtgttaaaa | aaggtgaagt | tttgggatta | ctaggacaca | atggagctgg | taaaagtact | 4620 |
| tccattaaaa | tgataactgg | gtgcacaaag | ccaactgcag | gagtgggtgt | gttacaaggc | 4680 |
| agcagagcat | cagtaaggca | acagcatgac | aacagcctca | agttcttggg | gtactgccct | 4740 |
| caggagaact | cactgtggcc | caagcttaca | atgaaagagc | acttggagtt | gtatgcagct | 4800 |
| gtgaaaggac | tgggcaaaga | agatgctgct | ctcagtattt | cacgatttgt | ggaagctctt | 4860 |
| aagctccagg | aacaacttaa | ggctcctgtg | aaaactctat | cagaggggaat | aaagagaaaag | 4920 |
| ctgtgctttg | tgctgagcat | cctggggaac | ccatcagtgg | tgcttctaga | tgagccgttc | 4980 |
| accgggatgg | accccgaggg | gcagcagcaa | atgtggcaga | tacttcaggc | taccgttaaa | 5040 |
| aacaaggaga | ggggcaccct | cttgaccacc | cattacatgt | cagaggctga | ggctgtgtgt | 5100 |
| gaccgatagg | ccatgatggg | gtcaggaacg | ctaaggtgta | ttggttccat | tcaacatctg | 5160 |
| aaaaacaagt | tttgtagaga | ttatttacta | gaaataaaaa | tgaaagaacc | taccaggtg | 5220 |
| gaagctctcc | acacagagat | tttgaagctt | ttccacagag | ctgcttggca | ggaaagatat | 5280 |
| tcctctttta | tggcgtataa | gttacctgtg | gaggatgtcc | accctctatc | tcgggccttt | 5340 |
| ttcaagttag | aggcgatgaa | acagaccttc | aacctggagg | aatacagcct | ctctcaggct | 5400 |
| accttggagc | aggtattctt | agaactctgt | aaagagcagg | agctgggaaa | tgttgatgat | 5460 |

| | | | | | | |
|-------------|-------------|-------------|-------------|------------|------------|------|
| aaaattgata | caacagttga | atggaaactt | ctcccacagg | aagaccctta | aaatgaagaa | 5520 |
| cctcctaaca | ttcaatttta | ggtcctacta | cattgttagt | ttccataatt | ctacaagaat | 5580 |
| gtttcctttt | acttcagtta | acaaaagaaa | acatttaata | aacattcaat | aatgattaca | 5640 |
| gttttcattt | ttaaaaattt | aggatgaagg | aaacaaggaa | atatagggaa | aagtagtaga | 5700 |
| caaaattaac | aaaatcagac | atgttattca | tccccaacat | gggtctattt | tgtgcttaaa | 5760 |
| aataatttaa | aatcataca | atattagggt | ggttttcggt | tattatcaat | aaagctaaca | 5820 |
| ctgagaacat | tttacaata | aaaatatgag | tttttttagcc | tgaacttcaa | atgtatcagc | 5880 |
| tatttttaaa | cattattttac | tcggattcta | atttaattgt | acattgacta | taagaaggtc | 5940 |
| tgataaaactg | atgaaatggc | acagcataac | atttaattat | aatgacattc | tgattataaa | 6000 |
| ataaatgcat | gtgaatttta | gtacatatgt | aagttatatg | gaagaagata | gccataatct | 6060 |
| gtaagaaagt | accgcagtta | atatttttctt | tagccaactt | atattcaatg | tattttttat | 6120 |
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<220>
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 35 40 45
 Ile Ser Ser Met Met His Pro Asn Lys Lys Tyr Glu Glu Val Pro Asn Ile
 50 55 60
 Glu Leu Asn Pro Met Asp Lys Phe Thr Leu Ser Asn Leu Ile Leu Gly
 65 70 75 80
 Tyr Thr Pro Val Thr Asn Ile Thr Ser Ser Ile Met Gln Lys Val Ser
 85 90 95
 Thr Asp His Leu Pro Asp Val Ile Ile Thr Glu Glu Tyr Thr Asn Glu
 100 105 110
 Lys Glu Met Leu Thr Ser Ser Leu Ser Lys Pro Ser Asn Phe Val Gly
 115 120 125
 Val Val Phe Lys Asp Ser Met Ser Tyr Glu Leu Arg Phe Phe Pro Asp
 130 135 140
 Met Ile Pro Val Ser Ser Ile Tyr Met Asp Ser Arg Ala Gly Cys Ser
 145 150 155 160
 Lys Ser Cys Glu Ala Ala Gln Tyr Trp Ser Ser Gly Phe Thr Val Leu
 165 170 175
 Gln Ala Ser Ile Asp Ala Ala Ile Ile Gln Leu Lys Thr Asn Val Ser
 180 185 190
 Leu Trp Lys Glu Leu Glu Ser Thr Lys Ala Val Ile Met Gly Glu Thr
 195 200 205

Ala Val Val Glu Ile Asp Thr Phe Pro Arg Gly Val Ile Leu Ile Tyr
 210 215 220
 Leu Val Ile Ala Phe Ser Pro Phe Gly Tyr Phe Leu Ala Ile His Ile
 225 230 235 240
 Val Ala Glu Lys Glu Lys Lys Ile Lys Glu Phe Leu Lys Ile Met Gly
 245 250 255
 Leu His Asp Thr Ala Phe Trp Leu Ser Trp Val Leu Leu Tyr Thr Ser
 260 265 270
 Leu Ile Phe Leu Met Ser Leu Leu Met Ala Val Ile Ala Thr Ala Ser
 275 280 285
 Leu Leu Phe Pro Gln Ser Ser Ser Ile Val Ile Phe Leu Leu Phe Phe
 290 295 300
 Leu Tyr Gly Leu Ser Ser Val Phe Phe Ala Leu Met Leu Thr Pro Leu
 305 310 315 320
 Phe Lys Lys Ser Lys His Val Gly Ile Val Glu Phe Phe Val Thr Val
 325 330 335
 Ala Phe Gly Phe Ile Gly Leu Met Ile Ile Leu Ile Glu Ser Phe Pro
 340 345 350
 Lys Ser Leu Val Trp Leu Phe Ser Pro Phe Cys His Cys Thr Phe Val
 355 360 365
 Ile Gly Ile Ala Gln Val Met His Leu Glu Asp Phe Asn Glu Gly Ala
 370 375 380
 Ser Phe Ser Asn Leu Thr Ala Gly Pro Tyr Pro Leu Ile Ile Thr Ile
 385 390 395 400
 Ile Met Leu Thr Leu Asn Ser Ile Phe Tyr Val Leu Leu Ala Val Tyr
 405 410 415
 Leu Asp Gln Val Ile Pro Gly Glu Phe Gly Leu Arg Arg Ser Ser Leu
 420 425 430
 Tyr Phe Leu Lys Pro Ser Tyr Trp Ser Lys Ser Lys Arg Asn Tyr Glu
 435 440 445
 Glu Leu Ser Glu Gly Asn Val Asn Gly Asn Ile Ser Phe Ser Glu Ile
 450 455 460
 Ile Glu Pro Val Ser Ser Glu Phe Val Gly Lys Glu Ala Ile Arg Ile
 465 470 475 480
 Ser Gly Ile Gln Lys Thr Tyr Arg Lys Lys Gly Glu Asn Val Glu Ala
 485 490 495
 Leu Arg Asn Leu Ser Phe Asp Ile Tyr Glu Gly Gln Ile Thr Ala Leu
 500 505 510
 Leu Gly His Ser Gly Thr Gly Lys Ser Thr Leu Met Asn Ile Leu Cys
 515 520 525
 Gly Leu Cys Pro Pro Ser Asp Gly Phe Ala Ser Ile Tyr Gly His Arg
 530 535 540
 Val Ser Glu Ile Asp Glu Met Phe Glu Ala Arg Lys Met Ile Gly Ile

| | | | | | | | | | | | | | | | | |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|-----|
| 545 | | | | | 550 | | | | | | 555 | | | | | 560 |
| Cys | Pro | Gln | Leu | Asp 565 | Ile | His | Phe | Asp | Val 570 | Leu | Thr | Val | Glu | Glu 575 | Asn | |
| Leu | Ser | Ile | Leu 580 | Ala | Ser | Ile | Lys | Gly 585 | Ile | Pro | Ala | Asn | Asn 590 | Ile | Ile | |
| Gln | Glu | Val 595 | Gln | Lys | Val | Leu | Leu 600 | Asp | Leu | Asp | Met | Gln 605 | Thr | Ile | Lys | |
| Asp | Asn 610 | Gln | Ala | Lys | Lys | Leu 615 | Ser | Gly | Gly | Gln | Lys 620 | Arg | Lys | Leu | Ser | |
| Leu 625 | Gly | Ile | Ala | Val | Leu 630 | Gly | Asn | Pro | Lys | Ile 635 | Leu | Leu | Leu | Asp | Glu 640 | |
| Pro | Thr | Ala | Gly | Met 645 | Asp | Pro | Cys | Ser | Arg 650 | His | Ile | Val | Trp | Asn 655 | Leu | |
| Leu | Lys | Tyr | Arg 660 | Lys | Ala | Asn | Arg | Val 665 | Thr | Val | Phe | Ser | Thr 670 | His | Phe | |
| Met | Asp | Glu 675 | Ala | Asp | Ile | Leu | Ala 680 | Asp | Arg | Lys | Ala | Val 685 | Ile | Ser | Gln | |
| Gly | Met 690 | Leu | Lys | Cys | Val | Gly 695 | Ser | Ser | Met | Phe | Leu 700 | Lys | Ser | Lys | Trp | |
| Gly 705 | Ile | Gly | Tyr | Arg | Leu 710 | Ser | Met | Tyr | Ile | Asp 715 | Lys | Tyr | Cys | Ala | Thr 720 | |
| Glu | Ser | Leu | Ser | Ser 725 | Leu | Val | Lys | Gln | His 730 | Ile | Pro | Gly | Ala | Thr 735 | Leu | |
| Leu | Gln | Gln | Asn 740 | Asp | Gln | Gln | Leu | Val 745 | Tyr | Ser | Leu | Pro | Phe 750 | Lys | Asp | |
| Met | Asp | Lys 755 | Phe | Ser | Gly | Leu | Phe 760 | Ser | Ala | Leu | Asp | Ser 765 | His | Ser | Asn | |
| Leu | Gly 770 | Val | Ile | Ser | Tyr | Gly 775 | Val | Ser | Met | Thr | Thr 780 | Leu | Glu | Asp | Val | |
| Phe 785 | Leu | Lys | Leu | Glu | Val 790 | Glu | Ala | Glu | Ile | Asp 795 | Gln | Ala | Asp | Tyr | Ser 800 | |
| Val | Phe | Thr | Gln | Gln 805 | Pro | Leu | Glu | Glu | Glu 810 | Met | Asp | Ser | Lys | Ser 815 | Phe | |
| Asp | Glu | Met | Glu 820 | Gln | Ser | Leu | Leu | Ile 825 | Leu | Ser | Glu | Thr | Lys 830 | Ala | Ser | |
| Leu | Val | Ser 835 | Thr | Met | Ser | Leu | Trp 840 | Lys | Gln | Gln | Met | Tyr 845 | Thr | Ile | Ala | |
| Lys | Phe 850 | His | Phe | Phe | Thr | Leu 855 | Lys | Arg | Glu | Ser | Lys 860 | Ser | Val | Arg | Ser | |
| Val 865 | Leu | Leu | Leu | Leu | Leu 870 | Ile | Phe | Phe | Thr | Val 875 | Gln | Ile | Phe | Met | Phe 880 | |
| Leu | Val | His | His | Ser 885 | Phe | Lys | Asn | Ala | Val 890 | Val | Pro | Ile | Lys | Leu 895 | Val | |

Pro Asp Leu Tyr Phe Leu Lys Pro Gly Asp Lys Pro His Lys Tyr Lys
 900 905 910
 Thr Ser Leu Leu Leu Gln Asn Ser Ala Asp Ser Asp Ile Ser Asp Leu
 915 920 925
 Ile Ser Phe Phe Thr Ser Gln Asn Ile Met Val Thr Met Ile Asn Asp
 930 935 940
 Ser Asp Tyr Val Ser Val Ala Pro His Ser Ala Ala Leu Asn Val Met
 945 950 955 960
 His Ser Glu Lys Asp Tyr Val Phe Ala Ala Val Phe Asn Ser Thr Met
 965 970 975
 Val Tyr Ser Leu Pro Ile Leu Val Asn Ile Ile Ser Asn Tyr Tyr Leu
 980 985 990
 Tyr His Leu Asn Val Thr Glu Thr Ile Gln Ile Trp Ser Thr Pro Phe
 995 1000 1005
 Phe Gln Glu Ile Thr Asp Ile Val Phe Lys Ile Glu Leu Tyr Phe Gln
 1010 1015 1020
 Ala Ala Leu Leu Gly Ile Ile Val Thr Ala Met Pro Pro Tyr Phe Ala
 1025 1030 1035 1040
 Met Glu Asn Ala Glu Asn His Lys Ile Lys Ala Tyr Thr Gln Leu Lys
 1045 1050 1055
 Leu Ser Gly Leu Leu Pro Ser Ala Tyr Trp Ile Gly Gln Ala Val Val
 1060 1065 1070
 Asp Ile Pro Leu Phe Phe Ile Ile Leu Ile Leu Met Leu Gly Ser Leu
 1075 1080 1085
 Leu Ala Phe His Tyr Gly Leu Tyr Phe Tyr Thr Val Lys Phe Leu Ala
 1090 1095 1100
 Val Val Phe Cys Leu Ile Gly Tyr Val Pro Ser Val Ile Leu Phe Thr
 1105 1110 1115 1120
 Tyr Ile Ala Ser Phe Thr Phe Lys Lys Ile Leu Asn Thr Lys Glu Phe
 1125 1130 1135
 Trp Ser Phe Ile Tyr Ser Val Ala Ala Leu Xaa Cys Ile Ala Ile Thr
 1140 1145 1150
 Glu Ile Thr Phe Phe Met Gly Tyr Thr Ile Ala Thr Ile Leu His Tyr
 1155 1160 1165
 Ala Phe Cys Ile Ile Ile Pro Ile Tyr Pro Leu Leu Gly Cys Leu Ile
 1170 1175 1180
 Ser Phe Ile Lys Ile Ser Trp Lys Asn Val Arg Lys Asn Val Asp Thr
 1185 1190 1195 1200
 Tyr Asn Pro Trp Asp Arg Leu Ser Val Ala Val Ile Ser Pro Tyr Leu
 1205 1210 1215
 Gln Cys Val Leu Trp Ile Phe Leu Leu Gln Tyr Tyr Glu Lys Lys Tyr
 1220 1225 1230

Gly Gly Arg Ser Ile Arg Lys Asp Pro Phe Phe Arg Asn Leu Ser Thr
 1235 1240 1245
 Lys Ser Lys Asn Arg Lys Leu Pro Glu Pro Pro Asp Asn Glu Asp Glu
 1250 1255 1260
 Asp Glu Asp Val Lys Ala Glu Arg Leu Lys Val Lys Glu Leu Met Gly
 1265 1270 1275 1280
 Cys Gln Cys Cys Glu Glu Lys Pro Ser Ile Met Val Ser Asn Leu His
 1285 1290 1295
 Lys Glu Tyr Asp Asp Lys Lys Asp Phe Leu Leu Ser Arg Lys Val Lys
 1300 1305 1310
 Lys Val Ala Thr Lys Tyr Ile Ser Phe Cys Val Lys Lys Gly Glu Ile
 1315 1320 1325
 Leu Gly Leu Leu Gly Pro Asn Gly Ala Gly Lys Ser Thr Ile Ile Asn
 1330 1335 1340
 Ile Leu Val Gly Asp Ile Glu Pro Thr Ser Gly Gln Val Phe Leu Gly
 1345 1350 1355 1360
 Asp Tyr Ser Ser Glu Thr Ser Glu Asp Asp Asp Ser Leu Lys Cys Met
 1365 1370 1375
 Gly Tyr Cys Pro Gln Ile Asn Pro Leu Trp Pro Asp Thr Thr Leu Gln
 1380 1385 1390
 Glu His Phe Glu Ile Tyr Gly Ala Val Lys Gly Met Ser Ala Ser Asp
 1395 1400 1405
 Met Lys Glu Val Ile Ser Arg Ile Thr His Ala Leu Asp Leu Lys Glu
 1410 1415 1420
 His Leu Gln Lys Thr Val Lys Lys Leu Pro Ala Gly Ile Lys Arg Lys
 1425 1430 1435 1440
 Leu Cys Phe Ala Leu Ser Met Leu Gly Asn Pro Gln Ile Thr Leu Leu
 1445 1450 1455
 Asp Glu Pro Ser Thr Gly Met Asp Pro Lys Ala Lys Gln His Met Trp
 1460 1465 1470
 Arg Ala Ile Arg Thr Ala Phe Lys Asn Arg Lys Arg Ala Ala Ile Leu
 1475 1480 1485
 Thr Thr His Tyr Met Glu Glu Ala Glu Ala Val Cys Asp Arg Val Ala
 1490 1495 1500
 Ile Met Val Ser Gly Gln Leu Arg Cys Ile Gly Thr Val Gln His Leu
 1505 1510 1515 1520
 Lys Ser Lys Phe Gly Lys Gly Tyr Phe Leu Glu Ile Lys Leu Lys Asp
 1525 1530 1535
 Trp Ile Glu Asn Leu Glu Val Asp Arg Leu Gln Arg Glu Ile Gln Tyr
 1540 1545 1550
 Ile Phe Pro Asn Ala Ser Arg Gln Glu Ser Phe Ser Ser Ile Leu Ala
 1555 1560 1565
 Tyr Lys Ile Pro Lys Glu Asp Val Gln Ser Leu Ser Gln Ser Phe Phe

1570 1575 1580
 Lys Leu Glu Glu Ala Lys His Ala Phe Ala Ile Glu Glu Tyr Ser Phe
 1585 1590 1595 1600
 Ser Gln Ala Thr Leu Glu Gln Val Phe Val Glu Leu Thr Lys Glu Gln
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 1635 1640

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 35 40 45
 Ser Ser Met Arg Asn Val Gln Phe Pro Gly Met Ala Pro Gln Asn Leu
 50 55 60
 Gly Arg Val Asp Lys Phe Asn Ser Ser Ser Leu Met Val Val Tyr Thr
 65 70 75 80
 Pro Ile Ser Asn Leu Thr Gln Gln Ile Met Asn Lys Thr Ala Leu Ala
 85 90 95
 Pro Leu Leu Lys Gly Thr Ser Val Ile Gly Ala Pro Asn Lys Thr His
 100 105 110
 Met Asp Glu Ile Leu Leu Glu Asn Leu Pro Tyr Ala Met Gly Ile Ile
 115 120 125
 Phe Asn Glu Thr Phe Ser Tyr Lys Leu Ile Phe Phe Gln Gly Tyr Asn
 130 135 140
 Ser Pro Leu Trp Lys Glu Asp Phe Ser Ala His Cys Trp Asp Gly Tyr
 145 150 155 160
 Gly Glu Phe Ser Cys Thr Leu Thr Lys Tyr Trp Asn Arg Gly Phe Val
 165 170 175
 Ala Leu Gln Thr Ala Ile Asn Thr Ala Ile Ile Glu Ile Thr Thr Asn
 180 185 190
 His Pro Val Met Glu Glu Leu Met Ser Val Thr Ala Ile Thr Met Lys
 195 200 205
 Thr Leu Pro Phe Ile Thr Lys Asn Leu Leu His Asn Glu Met Phe Ile
 210 215 220

Leu Phe Phe Leu Leu His Phe Ser Pro Leu Val Tyr Phe Ile Ser Leu
 225 230 235 240
 Asn Val Thr Lys Glu Arg Lys Lys Ser Lys Asn Leu Met Lys Met Met
 245 250 255
 Gly Leu Gln Asp Ser Ala Phe Trp Leu Ser Trp Gly Leu Ile Tyr Ala
 260 265 270
 Gly Phe Ile Phe Ile Ile Ser Ile Phe Ile Thr Ile Ile Ile Thr Phe
 275 280 285
 Thr Gln Ile Ile Val Met Thr Gly Phe Met Val Ile Phe Ile Leu Phe
 290 295 300
 Phe Leu Tyr Gly Leu Ser Leu Val Ala Leu Val Phe Leu Met Ser Val
 305 310 315 320
 Leu Leu Lys Lys Ala Val Leu Thr Asn Leu Val Val Phe Leu Leu Thr
 325 330 335
 Leu Phe Trp Gly Cys Leu Gly Phe Thr Val Phe Tyr Glu Gln Leu Pro
 340 345 350
 Ser Ser Leu Glu Trp Ile Leu Asn Ile Cys Ser Pro Phe Ala Phe Thr
 355 360 365
 Thr Gly Met Ile Gln Ile Ile Lys Leu Asp Tyr Asn Leu Asn Gly Val
 370 375 380
 Ile Phe Pro Asp Pro Ser Gly Asp Ser Tyr Thr Met Ile Ala Thr Phe
 385 390 395 400
 Ser Met Leu Leu Leu Asp Gly Leu Ile Tyr Leu Leu Leu Ala Leu Tyr
 405 410 415
 Phe Asp Lys Ile Leu Pro Tyr Gly Asp Glu Arg His Tyr Ser Pro Leu
 420 425 430
 Phe Phe Leu Asn Ser Ser Ser Cys Phe Gln His Gln Arg Thr Asn Ala
 435 440 445
 Lys Val Ile Glu Lys Glu Ile Asp Ala Glu His Pro Ser Asp Asp Tyr
 450 455 460
 Phe Glu Pro Val Ala Pro Glu Phe Gln Gly Lys Glu Ala Ile Arg Ile
 465 470 475 480
 Arg Asn Val Lys Lys Glu Tyr Lys Gly Lys Ser Gly Lys Val Glu Ala
 485 490 495
 Leu Lys Gly Leu Leu Phe Asp Ile Tyr Glu Gly Gln Ile Thr Ala Ile
 500 505 510
 Leu Gly His Ser Gly Ala Gly Lys Ser Ser Leu Leu Asn Ile Leu Asn
 515 520 525
 Gly Leu Ser Val Pro Thr Glu Gly Ser Val Thr Ile Tyr Asn Lys Asn
 530 535 540
 Leu Ser Glu Met Gln Asp Leu Glu Glu Ile Arg Lys Ile Thr Gly Val
 545 550 555 560
 Cys Pro Gln Phe Asn Val Gln Phe Asp Ile Leu Thr Val Lys Glu Asn

| 565 | | | | | | | | | | 570 | | | | | 575 | | | | |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|--|--|--|--|
| Leu | Ser | Leu | Phe 580 | Ala | Lys | Ile | Lys | Gly 585 | Ile | His | Leu | Lys | Glu 590 | Val | Glu | | | | |
| Gln | Glu | Val 595 | Gln | Arg | Ile | Leu | Leu 600 | Glu | Leu | Asp | Met | Gln 605 | Asn | Ile | Gln | | | | |
| Asp | Asn 610 | Leu | Ala | Lys | His | Leu 615 | Ser | Glu | Gly | Gln | Lys 620 | Arg | Lys | Leu | Thr | | | | |
| Phe 625 | Gly | Ile | Thr | Ile | Leu 630 | Gly | Asp | Pro | Gln | Ile 635 | Leu | Leu | Leu | Asp | Glu 640 | | | | |
| Pro | Thr | Thr | Gly | Leu 645 | Asp | Pro | Phe | Ser | Arg 650 | Asp | Gln | Val | Trp | Ser 655 | Leu | | | | |
| Leu | Arg | Glu | Arg 660 | Arg | Ala | Asp | His | Val 665 | Ile | Leu | Phe | Ser | Thr 670 | Gln | Ser | | | | |
| Met | Asp | Glu 675 | Ala | Asp | Ile | Leu | Ala 680 | Asp | Arg | Lys | Val | Ile 685 | Met | Ser | Asn | | | | |
| Gly | Arg 690 | Leu | Lys | Cys | Ala | Gly 695 | Ser | Ser | Met | Phe | Leu 700 | Lys | Arg | Arg | Trp | | | | |
| Gly 705 | Leu | Gly | Tyr | His | Leu 710 | Ser | Leu | His | Arg | Asn 715 | Glu | Ile | Cys | Asn | Pro 720 | | | | |
| Glu | Gln | Ile | Thr | Ser 725 | Phe | Ile | Thr | His | His 730 | Ile | Pro | Asp | Ala | Lys 735 | Leu | | | | |
| Lys | Thr | Glu | Asn 740 | Lys | Glu | Lys | Leu | Val 745 | Tyr | Thr | Leu | Pro | Leu 750 | Glu | Arg | | | | |
| Thr | Asn 755 | Thr | Phe | Pro | Asp | Leu | Phe 760 | Ser | Asp | Leu | Asp | Lys 765 | Cys | Ser | Asp | | | | |
| Gln | Gly 770 | Val | Thr | Gly | Tyr | Asp 775 | Ile | Ser | Met | Ser | Thr 780 | Leu | Asn | Glu | Val | | | | |
| Phe 785 | Met | Lys | Leu | Glu | Gly 790 | Gln | Ser | Thr | Ile | Glu 795 | Gln | Asp | Phe | Glu | Gln 800 | | | | |
| Val | Glu | Met | Ile | Arg 805 | Asp | Ser | Glu | Ser | Leu 810 | Asn | Glu | Met | Glu | Leu 815 | Ala | | | | |
| His | Ser | Ser | Phe 820 | Ser | Glu | Met | Gln | Thr 825 | Ala | Val | Ser | Asp | Met 830 | Gly | Leu | | | | |
| Trp | Arg | Met 835 | Gln | Val | Phe | Ala | Met 840 | Ala | Arg | Leu | Arg | Phe 845 | Leu | Lys | Leu | | | | |
| Lys | Arg 850 | Gln | Thr | Lys | Val | Leu 855 | Leu | Thr | Leu | Leu | Leu 860 | Val | Phe | Gly | Ile | | | | |
| Ala 865 | Ile | Phe | Pro | Leu | Ile 870 | Val | Glu | Asn | Ile | Ile 875 | Tyr | Ala | Met | Leu | Asn 880 | | | | |
| Glu | Lys | Ile | Asp | Trp 885 | Glu | Phe | Lys | Asn | Glu 890 | Leu | Tyr | Phe | Leu | Ser 895 | Pro | | | | |
| Gly | Gln | Leu | Pro 900 | Gln | Glu | Pro | Arg | Thr 905 | Ser | Leu | Leu | Ile | Ile 910 | Asn | Asn | | | | |

Thr Glu Ser Asn Ile Glu Asp Phe Ile Lys Ser Leu Lys His Gln Asn
 915 920 925
 Ile Leu Leu Glu Val Asp Asp Phe Glu Asn Arg Asn Gly Thr Asp Gly
 930 935 940
 Leu Ser Tyr Asn Gly Ala Ile Ile Val Ser Gly Lys Gln Lys Asp Tyr
 945 950 955 960
 Arg Phe Ser Val Val Cys Asn Thr Lys Arg Leu His Cys Phe Pro Ile
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 Leu Met Asn Ile Ile Ser Asn Gly Leu Leu Gln Met Phe Asn His Thr
 980 985 990
 Gln His Ile Arg Ile Glu Ser Ser Pro Phe Pro Leu Ser His Ile Gly
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 Leu Trp Thr Gly Leu Pro Asp Gly Ser Phe Phe Leu Phe Leu Val Leu
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 Cys Ser Ile Ser Pro Tyr Ile Thr Met Gly Ser Ile Ser Asp Tyr Lys
 1025 1030 1035 1040
 Lys Asn Ala Lys Ser Gln Leu Trp Ile Ser Gly Leu Tyr Thr Ser Ala
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 Arg Lys Arg Arg Lys Asn Ser Gly Leu Trp Ser Phe Tyr Phe Phe Phe
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 Ala Ser Thr Ile Met Phe Ser Ile Thr Leu Ile Asn His Phe Asp Leu
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 Ser Ile Leu Ile Thr Thr Met Val Leu Val Pro Ser Tyr Thr Leu Leu
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 Phe Pro Glu Ala Asn Phe Glu Leu Ser Ala Thr Asp Phe Leu Val Cys
 1185 1190 1195 1200
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Ile Asp Glu Asp Glu Asp Ile Gln Thr Glu Arg Ile Arg Thr Ala Thr
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 Val Lys Ser Ala Gln Ile Val Val Leu Thr Gly Phe Val Met Val Phe
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 Met Ser Val Leu Ile Lys Lys Pro Phe Leu Thr Gly Leu Val Val Phe
 325 330 335
 Leu Leu Ile Val Phe Trp Gly Ile Leu Gly Phe Pro Ala Leu Tyr Thr
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 His Leu Pro Ala Phe Leu Glu Trp Thr Leu Cys Leu Leu Ser Pro Phe
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 Ala Phe Thr Val Gly Met Ala Gln Leu Ile His Leu Asp Tyr Asp Val
 370 375 380
 Asn Ser Asn Ala His Leu Asp Ser Ser Gln Asn Pro Tyr Leu Ile Ile
 385 390 395 400
 Ala Thr Leu Phe Met Leu Val Phe Asp Thr Leu Leu Tyr Leu Val Leu
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 Thr Leu Tyr Phe Asp Lys Ile Leu Pro Ala Glu Tyr Gly His Arg Cys
 420 425 430
 Ser Pro Leu Phe Phe Leu Lys Ser Cys Phe Trp Phe Gln His Gly Arg
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 Ala Asn His Val Val Leu Glu Asn Glu Thr Asp Ser Asp Pro Thr Pro
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 Asn Asp Cys Phe Glu Pro Val Ser Pro Glu Phe Cys Gly Lys Glu Ala
 465 470 475 480
 Ile Arg Ile Lys Asn Leu Lys Lys Glu Tyr Ala Gly Lys Cys Glu Arg
 485 490 495
 Val Glu Ala Leu Lys Gly Val Val Phe Asp Ile Tyr Glu Gly Gln Ile
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 Thr Ala Leu Leu Gly His Ser Gly Ala Gly Lys Thr Thr Leu Leu Asn
 515 520 525
 Ile Leu Ser Gly Leu Ser Val Pro Thr Ser Gly Ser Val Thr Val Tyr
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 Asn His Thr Leu Ser Arg Met Ala Asp Ile Glu Asn Ile Ser Lys Phe
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 Thr Gly Phe Cys Pro Gln Ser Asn Val Gln Phe Gly Phe Leu Thr Val
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 Lys Glu Asn Leu Arg Leu Phe Ala Lys Ile Lys Gly Ile Leu Pro His
 580 585 590
 Glu Val Glu Lys Glu Val Gln Arg Val Val Gln Glu Leu Glu Met Glu

| 595 | | | | | 600 | | | | | 605 | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Asn | Ile | Gln | Asp | Ile | Leu | Ala | Gln | Asn | Leu | Ser | Gly | Gly | Gln | Asn | Arg |
| 610 | | | | | | 615 | | | | | 620 | | | | |
| Lys | Leu | Thr | Phe | Gly | Ile | Ala | Ile | Leu | Gly | Asp | Pro | Gln | Val | Leu | Leu |
| 625 | | | | | 630 | | | | | 635 | | | | | 640 |
| Leu | Asp | Glu | Pro | Thr | Ala | Gly | Leu | Asp | Pro | Leu | Ser | Arg | His | Arg | Ile |
| | | | | 645 | | | | | 650 | | | | | 655 | |
| Trp | Asn | Leu | Leu | Lys | Glu | Gly | Lys | Ser | Asp | Arg | Val | Ile | Leu | Phe | Ser |
| | | | 660 | | | | | 665 | | | | | 670 | | |
| Thr | Gln | Phe | Ile | Asp | Glu | Ala | Asp | Ile | Leu | Ala | Asp | Arg | Lys | Val | Phe |
| | | 675 | | | | | 680 | | | | | 685 | | | |
| Ile | Ser | Asn | Gly | Lys | Leu | Lys | Cys | Ala | Gly | Ser | Ser | Leu | Phe | Leu | Lys |
| | 690 | | | | | 695 | | | | | 700 | | | | |
| Lys | Lys | Trp | Gly | Ile | Gly | Tyr | His | Leu | Ser | Leu | His | Leu | Asn | Glu | Arg |
| 705 | | | | | 710 | | | | | 715 | | | | | 720 |
| Cys | Asp | Pro | Glu | Ser | Ile | Thr | Ser | Leu | Val | Lys | Gln | His | Ile | Ser | Asp |
| | | | | 725 | | | | | 730 | | | | | 735 | |
| Ala | Lys | Leu | Thr | Ala | Gln | Ser | Glu | Glu | Lys | Leu | Val | Tyr | Ile | Leu | Pro |
| | | | 740 | | | | | 745 | | | | | 750 | | |
| Leu | Glu | Arg | Thr | Asn | Lys | Phe | Pro | Glu | Leu | Tyr | Arg | Asp | Leu | Asp | Arg |
| | | 755 | | | | | 760 | | | | | 765 | | | |
| Cys | Ser | Asn | Gln | Gly | Ile | Glu | Asp | Tyr | Gly | Val | Ser | Ile | Thr | Thr | Leu |
| | 770 | | | | | 775 | | | | | 780 | | | | |
| Asn | Glu | Val | Phe | Leu | Lys | Leu | Glu | Gly | Lys | Ser | Thr | Ile | Asp | Glu | Ser |
| 785 | | | | | 790 | | | | | 795 | | | | | 800 |
| Asp | Ile | Gly | Ile | Trp | Gly | Gln | Leu | Gln | Thr | Asp | Gly | Ala | Lys | Asp | Ile |
| | | | | 805 | | | | | 810 | | | | | 815 | |
| Gly | Ser | Leu | Val | Glu | Leu | Glu | Gln | Val | Leu | Ser | Ser | Phe | His | Glu | Thr |
| | | | 820 | | | | | 825 | | | | | 830 | | |
| Arg | Lys | Thr | Ile | Ser | Gly | Val | Ala | Leu | Trp | Arg | Gln | Gln | Val | Cys | Ala |
| | | 835 | | | | | 840 | | | | | 845 | | | |
| Ile | Ala | Lys | Val | Arg | Phe | Leu | Lys | Leu | Lys | Lys | Glu | Arg | Lys | Ser | Leu |
| | 850 | | | | | 855 | | | | | 860 | | | | |
| Trp | Thr | Ile | Leu | Leu | Leu | Phe | Gly | Ile | Ser | Phe | Ile | Pro | Gln | Leu | Leu |
| 865 | | | | | 870 | | | | | 875 | | | | | 880 |
| Glu | His | Leu | Phe | Tyr | Glu | Ser | Tyr | Gln | Lys | Ser | Tyr | Pro | Trp | Glu | Leu |
| | | | | 885 | | | | | 890 | | | | | 895 | |
| Ser | Pro | Asn | Thr | Tyr | Phe | Leu | Ser | Pro | Gly | Gln | Gln | Pro | Gln | Asp | Pro |
| | | | 900 | | | | | 905 | | | | | 910 | | |
| Leu | Thr | His | Leu | Leu | Val | Ile | Asn | Lys | Thr | Gly | Ser | Thr | Ile | Asp | Asn |
| | | 915 | | | | | 920 | | | | | 925 | | | |
| Phe | Leu | His | Ser | Leu | Arg | Arg | Gln | Asn | Ile | Ala | Ile | Glu | Val | Asp | Ala |
| | 930 | | | | | 935 | | | | | 940 | | | | |

Phe Gly Thr Arg Asn Gly Thr Asp Asp Pro Ser Tyr Asn Gly Ala Ile
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Ile Val Ser Gly Asp Glu Lys Asp His Arg Phe Ser Ile Ala Cys Asn
965 970 975
Thr Lys Arg Leu Asn Cys Phe Pro Val Leu Leu Asp Val Ile Ser Asn
980 985 990
Gly Leu Leu Gly Ile Phe Asn Ser Ser Glu His Ile Gln Thr Asp Arg
995 1000 1005
Ser Thr Phe Phe Glu Glu His Met Asp Tyr Glu Tyr Gly Tyr Arg Ser
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Val Asp Val Ser Leu Tyr Phe Leu Ile Leu Leu Leu Met Gln Ile Met
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Asp Tyr Ile Phe Ser Pro Glu Glu Ile Ile Phe Ile Ile Gln Asn Leu
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Ala Thr Asp Leu Asn Glu Tyr Gly Phe Leu Gly Leu Phe Phe Gly Thr
1155 1160 1165
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1170 1175 1180
Glu Ile Ser Pro Asp Ser Met Asp Tyr Leu Gly Ala Ser Glu Ser Glu
1185 1190 1195 1200
Ile Val Tyr Leu Ala Leu Leu Ile Pro Tyr Leu His Phe Leu Ile Phe
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Leu Phe Ile Leu Arg Cys Leu Glu Met Asn Cys Arg Lys Lys Leu Met
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Arg Lys Asp Pro Val Phe Arg Ile Ser Pro Arg Ser Asn Ala Ile Phe
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Pro Asn Pro Glu Glu Pro Glu Gly Glu Glu Glu Asp Ile Gln Met Glu
1250 1255 1260
Arg Met Arg Thr Val Asn Ala Met Ala Val Arg Asp Phe Asp Glu Thr
1265 1270 1275 1280

Pro Val Ile Ile Ala Ser Cys Leu Arg Lys Glu Tyr Ala Gly Lys Lys
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 1345 1350 1355 1360
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 1365 1370 1375
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 Ile Lys Arg Lys Leu Arg Phe Val Leu Ser Ile Leu Gly Asn Pro Ser
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 1445 1450 1455
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 1475 1480 1485
 Asp Arg Val Ala Ile Met Val Ser Gly Arg Leu Arg Cys Ile Gly Ser
 1490 1495 1500
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<220>
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His Glu Met Val Gly Val Ile Phe Ser Asp Thr Phe Ser Tyr Arg Leu
      35      40      45
Lys Phe Asn Trp Gly Tyr Arg Ile Pro Val Ile Lys Glu His Ser Glu
      50      55      60
Tyr Thr Glu His Cys Trp Ala Met His Gly Glu Ile Phe Cys Tyr Leu
      65      70      75      80
Ala Lys Tyr Trp Leu Lys Gly Phe Val Ala Phe Gln Ala Ala Ile Asn
      85      90      95
Ala Ala Ile Ile Glu Val Thr Thr Asn His Ser Val Met Glu Glu Leu
      100      105
Thr Ser Val Ile Gly Ile Asn Met Lys Ile Pro Pro Phe Ile Ser Lys
      115      120      125
Gly Glu Ile Met Asn Glu Trp Phe His Phe Thr Cys Leu Val Ser Phe
      130      135      140
Ser Ser Phe Ile Tyr Phe Ala Ser Leu Asn Val Ala Arg Glu Arg Gly
      145      150      155      160
Lys Phe Lys Lys Leu Met Thr Val Met Gly Leu Arg Glu Ser Ala Phe
      165      170      175
Trp Leu Ser Trp Xaa Leu Thr Tyr Ile Cys Phe Ile Phe Ile Met Ser
      180      185      190
Ile Phe Met Ala Leu Val Ile Thr Ser Ile Ser Ile Val Phe His Thr
      195      200      205
Gly Phe Met Val Ile Phe Thr Leu Tyr Ser Leu Tyr Gly Leu Ser Leu
      210      215      220
Ile Ala Leu Ala Phe Leu Met Ser Val Leu Ile Arg Lys Pro Met Leu
      225      230      235      240
Ala Gly Leu Ala Gly Phe Leu Phe Thr Val Phe Trp Gly Cys Leu Gly
      245      250      255
Phe Thr Val Leu Tyr Arg Gln Leu Pro Leu Ser Leu Gly Trp Val Leu
      260      265      270

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Ser Leu Leu Ser Pro Phe Ala Phe Thr Ala Gly Met Ala Gln Val Thr
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 His Leu Asp Asn Tyr Leu Ser Gly Val Ile Phe Pro Asp Pro Ser Gly
 290 295 300
 Asp Ser Tyr Lys Met Ile Ala Thr Phe Phe Ile Leu Ala Phe Asp Thr
 305 310 315 320
 Leu Phe Tyr Leu Ile Phe Thr Leu Tyr Phe Glu Arg Val Leu Pro Asp
 325 330 335
 Lys Asp Gly His Gly Asp Ser Pro Leu Phe Phe Leu Lys Ser Ser Phe
 340 345 350
 Trp Ser Lys His Gln Asn Thr His His Glu Ile Phe Glu Asn Glu Ile
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 Asn Pro Glu His Ser Ser Asp Asp Ser Phe Glu Pro Val Ser Pro Glu
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 Phe His Gly Lys Glu Ala Ile Arg Ile Arg Asn Val Ile Lys Glu Tyr
 385 390 395 400
 Asn Gly Lys Thr Gly Lys Val Glu Ala Leu Gln Gly Ile Phe Phe Asp
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 Ile Tyr Glu Gly Gln Ile Thr Ala Ile Leu Gly His Asn Gly Ala Gly
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 Lys Ser Thr Leu Leu Asn Ile Leu Ser Gly Leu Ser Val Ser Thr Glu
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 Gly Ser Ala Thr Ile Tyr Asn Thr Gln Leu Ser Glu Ile Thr Asp Met
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 Glu Glu Ile Arg Lys Asn Ile Gly Phe Cys Pro Gln Phe Asn Phe Gln
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 Lys Gly Ile Gln Pro Lys Glu Val Glu Gln Glu Val Lys Arg Ile Ile
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 Met Glu Leu Asp Met Gln Ser Ile Gln Asp Ile Ile Ala Lys Lys Leu
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 Ser Gly Gly Gln Lys Arg Lys Leu Thr Leu Gly Ile Ala Ile Leu Gly
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 Asp Pro Gln Val Leu Leu Leu Asp Glu Pro Thr Ala Gly Leu Asp Pro
 545 550 555 560
 Phe Ser Arg His Arg Val Trp Ser Leu Leu Lys Glu His Lys Val Asp
 565 570 575
 Arg Leu Ile Leu Phe Ser Thr Gln Phe Met Asp Glu Ala Asp Ile Leu
 580 585 590
 Ala Asp Arg Lys Val Phe Leu Ser Asn Gly Lys Leu Lys Cys Ala Gly
 595 600 605

Ser Ser Leu Phe Leu Lys Arg Lys Trp Gly Ile Gly Tyr His Leu Ser
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 Leu His Arg Asn Glu Met Cys Asp Thr Glu Lys Ile Thr Ser Leu Ile
 625 630 635 640
 Lys Gln His Ile Pro Asp Ala Lys Leu Thr Thr Glu Ser Glu Glu Lys
 645 650 655
 Leu Val Tyr Ser Leu Pro Leu Glu Lys Thr Asn Lys Phe Pro Asp Leu
 660 665 670
 Tyr Ser Asp Leu Asp Lys Cys Ser Asp Gln Gly Ile Arg Asn Tyr Ala
 675 680 685
 Val Ser Val Thr Ser Leu Asn Glu Val Phe Leu Asn Leu Glu Gly Lys
 690 695 700
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 Thr His Cys Trp Glu Phe Ser Pro Ser Met Tyr Phe Leu Ser Leu Glu
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 Val Leu Glu Ile Asp Asp Phe Arg Asn Arg Asn Gly Ser Asp Asp Pro
 850 855 860
 Ser Tyr Asn Gly Ala Ile Ile Val Ser Gly Asp Gln Lys Asp Tyr Arg
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 Lys Arg Lys Leu Cys Phe Val Leu Ser Ile Leu Gly Asn Pro Ser Val
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 Tyr Lys Leu Pro Val Glu Asp Val His Pro Leu Ser Arg Ala Phe Phe
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 Ser Gln Ala Thr Leu Glu Gln Val Phe Leu Glu Leu Cys Lys Glu Gln
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<210> 16
 <211> 186
 <212> DNA
 <213> Homo sapiens

<400> 16
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 gtgtttctcc ttaccctctt ttggggatgt ctgggattca ctgtatttta tgaacaactt 120
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<210> 17
 <211> 148
 <212> DNA
 <213> Homo sapiens

<400> 17
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 gcattatact ttgacaaaat tttaccct 148

<210> 18
 <211> 169
 <212> DNA
 <213> Homo sapiens

<400> 18
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 accaaaggac taatgctaag gttattgaga aagaaatcga tgctgagcat ccctctgatg 120
 attattitga accagtagct cctgaattcc aaggaaaaga agccatcag 169

<210> 19
 <211> 59
 <212> DNA
 <213> Homo sapiens

<400> 19
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<210> 20
 <211> 111
 <212> DNA
 <213> Homo sapiens

<400> 20
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<210> 21
 <211> 176
 <212> DNA
 <213> Homo sapiens

<400> 21
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 agataactgg cgtctgtcct caattcaatg ttcaatttga catactcacc gtgaaggaaa 120
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<210> 22
 <211> 120
 <212> DNA
 <213> Homo sapiens

<400> 22
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 ttaagtgaag gacagaaaag aaagctgact tttgggatta ccattttagg agatcctcaa 120

<210> 23
 <211> 139
 <212> DNA
 <213> Homo sapiens

<400> 23
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 gaggctgaca tcctggctg 139

<210> 24
 <211> 91
 <212> DNA
 <213> Homo sapiens

<400> 24
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 aaagaagggt gggctcttga taccaccta g 91

<210> 25
 <211> 140
 <212> DNA
 <213> Homo sapiens

<400> 25
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 aaggacaaat acatttcag 140

<210> 26
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 <212> DNA
 <213> Homo sapiens

<400> 26
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 tgtcaactct aaatgaagtc tttatgaaac tggaaggaca gtcaactatc gaacaag 117

<210> 27
 <211> 184
 <212> DNA
 <213> Homo sapiens

<400> 27
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 ccct 184

<210> 28
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<212> DNA
<213> Homo sapiens

<400> 28
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gttaaattgaa aagatcgatt gggaatttaa aaacgaattg tattttctct ctcctggaca 120
acttccccag gaaccccgta ccagcctggt gatcatcaat aacacag 167

<210> 29
<211> 134
<212> DNA
<213> Homo sapiens

<400> 29
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atgactttga aaacagaaat ggtactgatg gcctctcata caatggagct atcatagttt 120
ctggtaaaca aaag 134

<210> 30
<211> 138
<212> DNA
<213> Homo sapiens

<400> 30
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aatattatca gcaatgggct acttcaaagt tttaatcaca cacaacatat tcgaattgag 120
tcaagcccat ttcctctt 138

<210> 31
<211> 108
<212> DNA
<213> Homo sapiens

<400> 31
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tgtagcattt ctccttatat caccatgggc agcatcagtg attacaag 108

<210> 32
<211> 174
<212> DNA
<213> Homo sapiens

<400> 32
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caggcactag tggacgtcag cttcttcatt ttaattctcc ttttaatgta ttttaatttc 120
tacatagaaa acatgcagta cttctttatt acaagccaaa ttgtgtttgc tttg 174

<210> 33
<211> 114
<212> DNA
<213> Homo sapiens

<400> 33
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<210> 34
<211> 120
<212> DNA
<213> Homo sapiens

<400> 34
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accaccatgg tattgggtcc ttcatatacc ttgcttgat ttaaaacttt tttggaagtg 120

<210> 35
<211> 78
<212> DNA
<213> Homo sapiens

<400> 35
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tttctagtct gcttcata 78

<210> 36
<211> 92
<212> DNA
<213> Homo sapiens

<400> 36
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aagaaaagaa tgcgaaaaga tcctgttttc ag 92

<210> 37
<211> 121
<212> DNA
<213> Homo sapiens

<400> 37
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agatattcaa acagaaagaa taagaacagc cactgctctg accacttcaa tcttagatga 120
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<210> 38
<211> 118
<212> DNA
<213> Homo sapiens

<400> 38
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tttcaaaga ggaagaagaa aatagcagca agaaatatct ctttctgtgt tcaagaag 118

<210> 39
<211> 92
<212> DNA
<213> Homo sapiens

<400> 39
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tatctgggat caciaagcca actgctggag ag 92

<210> 40
<211> 155

<212> DNA
<213> Homo sapiens

<400> 40
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gtgctgtggc ccatgctgac gttgagggaa cacctggagg tgtatgctgc cgtcaagggg 120
ctcaggaaag cggacgcgag gctcgccatc gcaag 155

<210> 41
<211> 76
<212> DNA
<213> Homo sapiens

<400> 41
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aggaatcacg agaaag 76

<210> 42
<211> 95
<212> DNA
<213> Homo sapiens

<400> 42
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acgggcatag accccacagg gcagcagcaa atgtg 95

<210> 43
<211> 120
<212> DNA
<213> Homo sapiens

<400> 43
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cctggctgag gcggaagcct tgtgtgaccg tgtggccatc atgggtgtctg gaaggcttag 120

<210> 44
<211> 141
<212> DNA
<213> Homo sapiens

<400> 44
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aaaagtgaag gaaacgtctc aagtgacttt ggtccacact gagattctga agcttttccc 120
acaggctgca gggcaggaaa g 141

<210> 45
<211> 80
<212> DNA
<213> Homo sapiens

<400> 45
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ctttcacaaa ttagaagcag 80

<210> 46
<211> 56
<212> DNA
<213> Homo sapiens

<400> 46
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<210> 47
<211> 369
<212> DNA
<213> Homo sapiens

<400> 47
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acaatgagat ggaaactcct ccctcattca gatgaacctt aaaacctcaa acctagtaat 120
tttttggtga tctcctataa acttatgttt tatgtaataa ttaatagtat gtttaatttt 180
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ttttaaaatt attcttcctc tcaaacatag ggggtgatagc aaacctgtga taaaggcaat 300
acaaaatatt agtaaagtca cccaaagagt caggcactgg gtattgtgga aataaaaacta 360
tataaactt 369

<210> 48
<211> 130
<212> DNA
<213> Homo sapiens

<400> 48
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tctcctccag aacatgcaga gacccatgga tgaactgtgt ttctagattt ttcctccagc 120
ttcctgaga 130

<210> 49
<211> 109
<212> DNA
<213> Homo sapiens

<400> 49
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ctgcaagaac tgtctcaaaa aatggagaat gaaaagacag accttggtg 109

<210> 50
<211> 208
<212> DNA
<213> Homo sapiens

<400> 50
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caagttcatg acactcctca aatgtcttca atggatctgg gacgtgtaga tagttttaat 120
gatactaatt atgttattgc atttgcacct gaatccaaaa ctaccaaga gataatgaac 180
aaagtggctt cagccccatt cctaaaag 208

<210> 51
<211> 165
<212> DNA
<213> Homo sapiens

<400> 51
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caatagacgc agtgagagtc atctttactg ataccttctc ctaccatttg aagttttctt 120
ggggacatag aatccccatg atgaaagagc acagagacca ttcag 165

<210> 52
<211> 104
<212> DNA
<213> Homo sapiens

<400> 52
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gctttgtagc ttttcaagct gccattaatg ctgctatcat agaa 104

<210> 53
<211> 227
<212> DNA
<213> Homo sapiens

<400> 53
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atattacctt ttgttgccca aggaggagtt gcaactgatt ttttcatttt cttttgcatt 120
atttcctttt ctacatttat atactatgta tcagtcaatg ttacacaaga aagacaatac 180
attacgtcat tgatgacaat gatgggactc cgagagtcag cattctg 227

<210> 54
<211> 142
<212> DNA
<213> Homo sapiens

<400> 54
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tctcctctat ggcctgtcct tg 142

<210> 55
<211> 186
<212> DNA
<213> Homo sapiens

<400> 55
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gtgtttctcc ttattgtcct ttgggggatc ctgggattcc cagcattgta tacacatctt 120
cctgcatttt tggaatggac tttgtgtcct cttagcccct ttgccttcac tgttgggatg 180
gccag 186

<210> 56
<211> 148
<212> DNA
<213> Homo sapiens

<400> 56
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tacctcataa tagctactct tttcatgttg gtttttgaca cccttctgta tttggtattg 120
acattatatt ttgacaaaat tttgcccg 148

<210> 57
<211> 169
<212> DNA
<213> Homo sapiens

<400> 57
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acggaagggc taatcatgtg gtccttgaga atgaaacaga ttctgatcct acacctaattg 120

actgttttga accagtgtct ccagaattct gtgggaagga agccatcag 169

<210> 58
<211> 59
<212> DNA
<213> Homo sapiens

<400> 58
aatcaaaaat cttaaaaaag aatatgcagg gaagtgtgag agagtagaag ctttgaaag 59

<210> 59
<211> 111
<212> DNA
<213> Homo sapiens

<400> 59
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<210> 60
<211> 176
<212> DNA
<213> Homo sapiens

<400> 60
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agttcactgg attttgtcca caatccaatg tgcaatttgg atttctcact gtgaaagaaa 120
acctcaggct gtttgctaaa ataaaaggga ttttgccaca tgaagtggag aaagag 176

<210> 61
<211> 120
<212> DNA
<213> Homo sapiens

<400> 61
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ttaagtgggt gacaaaatag gaaactaact tttgggattg ccattttagg agatcctcaa 120

<210> 62
<211> 139
<212> DNA
<213> Homo sapiens

<400> 62
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aatctcctga aagaggggaa atcagacaga gtaattctct tcagcaccca gtttatagat 120
gaggctgaca ttctggcgg 139

<210> 63
<211> 91
<212> DNA
<213> Homo sapiens

<400> 63
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agaagaaatg gggcataggc taccatttaa g 91

<210> 64
 <211> 140
 <212> DNA
 <213> Homo sapiens

<400> 64
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 aaggacaaac aaatttccag 140

<210> 65
 <211> 120
 <212> DNA
 <213> Homo sapiens

<400> 65
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 taacaacttt gaatgaggtg tttctgaaat tagaaggaaa atcaactatt gatgaatcag 120

<210> 66
 <211> 199
 <212> DNA
 <213> Homo sapiens

<400> 66
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 tctggaggca gcaggctctgt gcaatagcaa aagttcgctt cctaaagtta aagaaagaaa 180
 gaaaaagcct gtggactat 199

<210> 67
 <211> 167
 <212> DNA
 <213> Homo sapiens

<400> 67
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 atatcagaaa agttaccctg gggaactgtc tccaaatata tacttcctct caccaggaca 120
 acaaccacag gatcctctga cccatttact ggtcatcaat aagacag 167

<210> 68
 <211> 134
 <212> DNA
 <213> Homo sapiens

<400> 68
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 caggtgatga aaag 134

<210> 69
 <211> 138
 <212> DNA
 <213> Homo sapiens

<400> 69
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 gatgtcatta gcaatggact acttgaattt ttttaattcgt cagaacacat tcagactgac 120
 agaagcacat tttttgaa 138

<210> 70
<211> 108
<212> DNA
<213> Homo sapiens

<400> 70
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gcctctttca ctccatacat tgcaatgagc agcattggtg actacaaa 108

<210> 71
<211> 174
<212> DNA
<213> Homo sapiens

<400> 71
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caagcactgg tggatgtttc cctgtacttt ttgatcctcc tgctaattgca aataatggat 120
tatattttta gcccagagga gattatattt ataattcaaa acctgttaat tcaa 174

<210> 72
<211> 114
<212> DNA
<213> Homo sapiens

<400> 72
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<210> 73
<211> 120
<212> DNA
<213> Homo sapiens

<400> 73
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<210> 74
<211> 69
<212> DNA
<213> Homo sapiens

<400> 74
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<210> 75
<211> 92
<212> DNA
<213> Homo sapiens

<400> 75
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aagaaactaa tgagaaagga tcctgtgttc ag 92

<210> 76

<211> 121
<212> DNA
<213> Homo sapiens

<400> 76
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<210> 77
<211> 118
<212> DNA
<213> Homo sapiens

<400> 77
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<210> 78
<211> 92
<212> DNA
<213> Homo sapiens

<400> 78
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<210> 79
<211> 161
<212> DNA
<213> Homo sapiens

<400> 79
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<210> 80
<211> 76
<212> DNA
<213> Homo sapiens

<400> 80
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gggaataaag cgaaag 76

<210> 81
<211> 95
<212> DNA
<213> Homo sapiens

<400> 81
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accgggatgg accccgaggg gcagcagcaa atgtg 95

<210> 82
<211> 120
<212> DNA

<213> Homo sapiens

<400> 82

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<210> 83

<211> 141

<212> DNA

<213> Homo sapiens

<400> 83

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ccaggctgct cagcaggaaa g 141

<210> 84

<211> 80

<212> DNA

<213> Homo sapiens

<400> 84

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tttcttcaaa ttagagatag 80

<210> 85

<211> 56

<212> DNA

<213> Homo sapiens

<400> 85

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<210> 86

<211> 1062

<212> DNA

<213> Homo sapiens

<400> 86

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<211> 287
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 <213> Homo sapiens

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 taacatgtaa ataggcatta atttttgaga aatagaaatg tttatcctta atgtattttt 180
 aatttgctaa cattgatttt ttatttttctt tcctgaaata gcttatttcc taaaatgaaa 240
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 cacagaccat ggcttttcag aagccaagct gaataaaaaac agtttttaaaa gaggcaacca 180
 ttgttagagg agtccttgaa ggattcttca ttgttttctt ggacaaaaag agaccagtgg 240
 atccaagtgc ttcaaatact tctctcttat tttcttaact 280

<210> 89
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 <212> DNA
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<400> 89
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<210> 91
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 ggggatatag aatcccagtt ataaaggagc actctgaata cacag 165

<210> 92
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 <212> DNA
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<210> 93
<211> 227
<212> DNA
<213> Homo sapiens

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ataccacctt tcattttctaa gggagaaatt atgaatgaat ggtttcattt tacttgctta 120
gtttctttct cttcttttat atactttgca tcattaaatg ttgcaaggga aagaggaaaa 180
tttaagaaac tgatgacagt aatgggtctc cgagagtcag cattctg 227

<210> 94
<211> 142
<212> DNA
<213> Homo sapiens

<400> 94
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ggtcataaca tcaatctcaa ttgtatttca tactggcttc atgggtgatat tcacactcta 120
tagcttatat ggcctttctt tg 142

<210> 95
<211> 186
<212> DNA
<213> Homo sapiens

<400> 95
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ggatttctct tcactgtatt ttggggatgt ctgggattca ctgtgttata cagacaactt 120
cctttatctt tgggatgggt attaatctt cttagccctt ttgccttcac tgctggaatg 180
gcccag 186

<210> 96
<211> 148
<212> DNA
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<400> 96
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acattatatt ttgagcgagt tttacctg 148

<210> 97
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<212> DNA
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attcttttga accggtgtct ccagaattcc atggaaaaga agccataag 169

<210> 98
<211> 59

<212> DNA
<213> Homo sapiens

<400> 98
aatcagaaat gttataaaaag aatataatgg aaagactgga aaagtagaag cattgcaag 59

<210> 99
<211> 111
<212> DNA
<213> Homo sapiens

<400> 99
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gtaaatcaac actgctaaac attcttagtg gattgtctgt ttctacagaa g 111

<210> 100
<211> 176
<212> DNA
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<210> 101
<211> 120
<212> DNA
<213> Homo sapiens

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<210> 102
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<212> DNA
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<400> 107
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<210> 108
 <211> 134
 <212> DNA
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<400> 108
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 <211> 138
 <212> DNA
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<213> Homo sapiens

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caggctctgg tggacattcc attatacttc ttgattctct tttcaatata ttttaatttac 120
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<210> 112

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<212> DNA

<213> Homo sapiens

<400> 112

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<210> 114

<211> 81

<212> DNA

<213> Homo sapiens

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<210> 115

<211> 92

<212> DNA

<213> Homo sapiens

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<213> Homo sapiens

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<400> 118
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<400> 119
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 <212> DNA
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| <400> 165 | |
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